

**Amendments to the Claims:**

This listing of claims will replace all prior versions, and listings, of claims in the application:

**Listing of Claims:**

1. (Currently amended) A polypeptide variant with increased heparin-binding ability, characterized in that
  - (i) added to the amino acid sequence of a polypeptide is at least one oligopeptide comprising the amino acid sequence  $X_1X_2X_3X_4X_5X_6$ ; and/or
  - (ii) inserted into the amino acid sequence of a polypeptide is at least one oligopeptide comprising the amino acid sequence  $X_1X_2X_3X_4X_5X_6$ ; and/or
  - (iii) at least one oligopeptide sequence naturally occurring within the amino acid sequence of a polypeptide is substituted by an oligopeptide comprising an amino acid sequence  $X_1X_2X_3X_4X_5X_6$ ,

wherein

$X_1 = K, R, \text{ or } H$ ;  
 $X_2 = K, R, \text{ or } H$ ;  
 $X_3 = K, R, H, \text{ or no amino acid}$ ;  
 $X_4 = \text{not } K, R, H, \text{ but any other amino acid}$ ;  
 $X_5 = \text{not } K, R, H, \text{ but any other or no amino acid}$ ;  
 $X_6 = \text{not } K, R, H, \text{ but any other or no amino acid (SEQ ID NO: 1)}$ ,

or

~~$X_1 = K, R, \text{ or } H$ ;~~  
 ~~$X_2 = \text{not } K, R, H, \text{ but any other amino acid}$ ;~~  
 ~~$X_3 = K, R, \text{ or } H$ ;~~  
 ~~$X_4 = \text{not } K, R, H, \text{ but any other amino acid}$ ;~~  
 ~~$X_5 = \text{not } K, R, H, \text{ but any other or no amino acid}$ ;~~  
 ~~$X_6 = \text{not } K, R, H, \text{ but any other or no amino acid (SEQ ID NO: 2)}$ ; and~~

the polypeptide is ~~selected from among members of the DVR family including the TGF- $\beta$  superfamily~~ a bone morphogenetic protein (BMP) or a growth differentiation factor (GDF).

2. (Original) A polypeptide variant as recited in claim 1, characterized in that one to four copies of said oligopeptide are inserted at one to four positions within the polypeptide.

3. (Previously presented) A polypeptide variant as recited in claim 1, characterized in said oligopeptide comprises amino acid sequence RKRA (SEQ ID NO:3) or RKRAKHKQ (SEQ ID NO:4).

4. (Previously presented) A polypeptide variant as recited in claim 1, characterized in said oligopeptide is added to the N-terminus and/or inserted into the N-terminal region, and/or substitutes a part of the N-terminal region.

5. (Previously presented) A polypeptide variant as recited in claim 1, characterized in that the amino acid sequence of said polypeptide variant further contains a sequence of relevance to recombinant expression at the N-terminus, said sequence of relevance to recombinant expression being M or MZ, where M stands for methionine and Z stands for one or more amino acids.

6. (Previously presented) A polypeptide variant as recited in claim 1, characterized in that said polypeptide variant further contains a His-tag.

7. (Previously presented) A polypeptide variant as recited in claim 1, characterized in that said polypeptide is altered by addition, substitution, insertion, inversion, and/or deletion, where said polypeptide altered by addition, substitution, insertion, inversion and/or deletion shows at least 10% of the biological activity of the unaltered polypeptide, and at least 90% homology to the unaltered polypeptide.

8. (Currently amended) A polypeptide variant as recited in claim 1, characterized in that said polypeptide is BMP-2, BMP-4, BMP-5, BMP-6, BMP-7/OP-1, or BMP-8/OP-2, or GDF-5.

9. (Previously presented) A polypeptide variant as recited in claim 1, wherein the polypeptide has a cysteine knot, characterized in that said oligopeptide is inserted before the cysteine knot.

10. (Previously presented) A polypeptide variant as recited in claim 8, characterized in that said polypeptide variant has the amino acid sequence SEQ ID NO:5 (T3) or SEQ ID NO:6 (T4).

11. (Previously presented) A polypeptide variant as recited in claim 1, characterized in that said polypeptide variant is a polymer, oligomer, or dimer.

12. (Previously presented) A nucleic acid molecule, comprising a nucleic acid sequence encoding a polypeptide variant as recited in claim 1.

13. (Original) A nucleic acid molecule as recited in claim 12, characterized in that said nucleic acid sequence is derived from genomic DNA or cDNA, or is a synthetic DNA.

14. (Previously presented) A nucleic acid molecule as recited in claim 12, further comprising a promoter suited to control expression, wherein said nucleic acid sequence encoding a polypeptide variant is under the control of said promoter.

15. (Previously presented) A nucleic acid molecule as recited in claim 12, wherein said nucleic acid molecule contains at least part of a vector.

16. (Previously presented) An isolated host cell, containing a nucleic acid molecule as recited in claim 12, wherein said host cell is a prokaryotic or eukaryotic cell suitable for expression of said nucleic acid molecule.

17. (Currently amended) A process for producing a polypeptide variant with increased heparin-binding ability as recited in claim 1, comprising:

addition to the amino acid sequence of a polypeptide of at least one oligopeptide containing an amino acid sequence selected from SEQ ID NO:1 ~~or SEQ ID NO:2~~; and/or

insertion into the amino acid sequence of a polypeptide of at least one oligopeptide containing an amino acid sequence selected from SEQ ID NO:1 ~~or SEQ ID NO:2~~; and/or

substitution of at least one oligopeptide sequence naturally occurring within the amino acid sequence of a polypeptide by one oligopeptide containing an amino acid sequence selected from SEQ ID NO:1 ~~or SEQ ID NO:2~~.

18. (Original) A process as recited in claim 17, characterized in that said process comprises a chemical and/or enzymatic synthesis process.

19. (Previously presented) A process as recited in claim 17, characterized in that said process comprises gene technological processes.

20. (Currently amended) A process as recited in claim 17, characterized in that said process comprises:

a) in vitro mutagenesis of a nucleic acid encoding a polypeptide, so that  
(i) to the nucleic acid encoding said polypeptide is added at least one nucleic acid encoding an oligopeptide containing an amino acid sequence that is selected from SEQ ID NO:1 ~~or SEQ ID NO:2~~; and/or

(ii) into the nucleic acid encoding said polypeptide is inserted at least one nucleic acid encoding an oligopeptide containing an amino acid sequence that is selected from SEQ ID NO:1 ~~or SEQ ID NO:2~~; and/or

(iii) at least one nucleic acid sequence naturally occurring within the nucleic acid sequence encoding said polypeptide is substituted by a nucleic acid sequence encoding an oligopeptide containing an amino acid sequence selected from SEQ ID NO:1 ~~or SEQ ID NO:2~~;

b) cloning of the mutated nucleic acid into a suitable expression vector;

- c) transformation/transfection of a suitable host cell with the expression vector obtained;
- d) cultivation of said transformed/transfected host cell under conditions suitable for expression;
- e) isolation, and if necessary renaturation, of the expressed polypeptide variant.

21. (Previously presented) A process as recited in claim 17, characterized in that said process is carried out within a prokaryotic host cell.

22. (Previously presented) A process as recited in claim 17, characterized in that said process is carried out within a eukaryotic cell.

23. (Previously presented) A pharmaceutical composition, comprising a polypeptide variant as recited in claim 1.

24. (Canceled)

25. (Currently amended) A composition for osteoinduction, comprising a polypeptide variant as recited in claim 1 and a carrier selected from among heparin, hydroxyapatite, hyaluronic acid, synthetic polymers, and collagen.

26. (Currently amended) ~~An osteoinductive~~ A matrix, characterized in that said matrix contains or is coated with heparin or heparin-like substances and polypeptide variants as recited in claim 1 are adsorbed to said heparin or heparin-like substances.

27. (Previously presented) A process as recited in claim 21, wherein the prokaryotic host cell is *E. coli*.

28. (Previously presented) A process as recited in claim 22, wherein the eukaryotic cell is selected from the group consisting of a yeast cell, a plant cell, an insect cell, CHO cells, and COS cells.

29. (Previously presented) A pharmaceutical composition as recited in claim 23, further comprising physiologically compatible additives.